

Persyn, a Member of the Synuclein Family, Has a Distinct Pattern of Expression in the Developing Nervous System

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The synucleins are a unique family of small intracellular proteins that have recently attracted considerable attention because of their involvement in human neurodegenerative diseases. We have cloned a new member of the synuclein family called persyn. In contrast to other synucleins, which are presynaptic proteins of CNS neurons, persyn is a cytosolic protein that is expressed predominantly in the cell bodies and axons of primary sensory neurons, sympathetic neurons, and motoneurons. Northern blotting, *in situ* hybridization, Western blotting,

and immunohistochemistry revealed that persyn mRNA and protein are expressed in these neurons from the earliest stages of axonal outgrowth and are maintained at a high level throughout life. Persyn also becomes detectable in evolutionary recent regions of the brain by adulthood.

Key words: synucleins; neurodegenerative diseases; Alzheimer's disease; Parkinson's disease; development of the nervous system; BCSG1; motoneurons; sensory neurons

The accumulation of intracellular or extracellular deposits in the form of plaques, tangles, or inclusion bodies is a hallmark of many neurodegenerative diseases (Selkoe, 1994; Kelly, 1996). These deposits are composed of one or more proteins and peptides, and it has been suggested that certain minor polypeptide components initiate aggregation of the main component (Wisniewski and Frangione, 1992; Ma et al., 1994). One of such minor components found in the plaques of Alzheimer's disease is NAC, an internal peptide of NACP or α -synuclein, a member of a family of small proteins with an unusual amino acid sequence and undefined functions (Maroteaux et al., 1988; Maroteaux and Scheller, 1991; Uéda et al., 1993; Jakes et al., 1994). The ability of α -synuclein to self-aggregate and to bind and induce the aggregation of amyloid A β peptide (Iwai et al., 1995b; Yoshimoto et al., 1995; Jensen et al., 1997; Paik et al., 1997, 1998) is consistent with a potential role in promoting amyloid deposition in Alzheimer's disease. The topography of synaptic function impairment in Alzheimer's disease also correlates with the pattern of α -synuclein expression and its localization in synaptic terminals (Uéda et al., 1993; Jakes et al., 1994; George et al., 1995; Iwai et al., 1995a). Furthermore, synuclein (synelfin) has been implicated in at least one form of learning and memory in zebrafish (George et al., 1995; Jin and Clayton, 1997). Both α - and β -synucleins are expressed predominantly in evolutionary recent structures of the CNS (Maroteaux and Scheller, 1991; Uéda et al., 1993, 1994; Jakes et al., 1994).

Two mutations in the α -synuclein gene have recently been reported to be directly associated with an early-onset, autosomal-dominant form of Parkinson's disease (Polymeropoulos et al., 1997; Kruger et al., 1998). Although it has been shown that

synucleins in solution are random coiled or natively unfolded proteins, interactions with other macromolecules in living cells could stabilize synucleins in particular conformations (Weinreb et al., 1996; Kim, 1997; Paik et al., 1997, 1998; Davidson et al., 1998). Both mutations associated with Parkinson's disease were predicted to change the secondary structure of the α -synuclein molecule in the way that increases the ability of this protein to self-aggregate or form aggregates with other proteins (Goedert, 1997; Heintz and Zoghbi, 1997; Nussbaum and Polymeropoulos, 1997; Polymeropoulos et al., 1997; Kruger et al., 1998). Consistently, α -synuclein, but not β -synuclein, is found in the Lewy bodies of cases of sporadic Parkinson's disease and dementia with Lewy bodies (Spillantini et al., 1997). Accumulations of α -synuclein are also present in abnormal neurites that, like Lewy bodies, contain ubiquitin, synaptophysin, and neurofilaments (Takeda et al., 1998). In contrast, τ -positive lesions (neurofibrillary tangles, Pick's bodies, neuropil threads, and ballooned neurons) are synuclein-negative (Takeda et al., 1998). These data suggest that α -synuclein is involved in specific aspects of the pathogenesis of neurodegeneration.

Here we report the characterization a member of the synuclein family that we have called persyn. Detailed studies of the expression of *persyn* mRNA and protein during normal embryonic and postnatal development show that it has a very different pattern of expression to that of other synucleins. This finding together with our evidence implicating persyn in regulating the integrity of the neurofilament network (Buchman et al., 1998) suggests that persyn could be involved in modulating axonal architecture during development and in the adult.

MATERIALS AND METHODS

Subtractive cloning. A subtractive cloning procedure (Baka and Buchman, 1996) was used to isolate cDNAs corresponding to mRNAs that are expressed at higher levels in embryonic day (E13) mouse trigeminal ganglia (TG) than in E13 mouse forebrain. Single-stranded cDNA was synthesized from E13 TG poly(A)⁺ RNA (leader) by reverse transcrip-

Received June 19, 1998; revised Aug. 18, 1998; accepted Aug. 28, 1998.

This work was supported by a grant from the Wellcome Trust.

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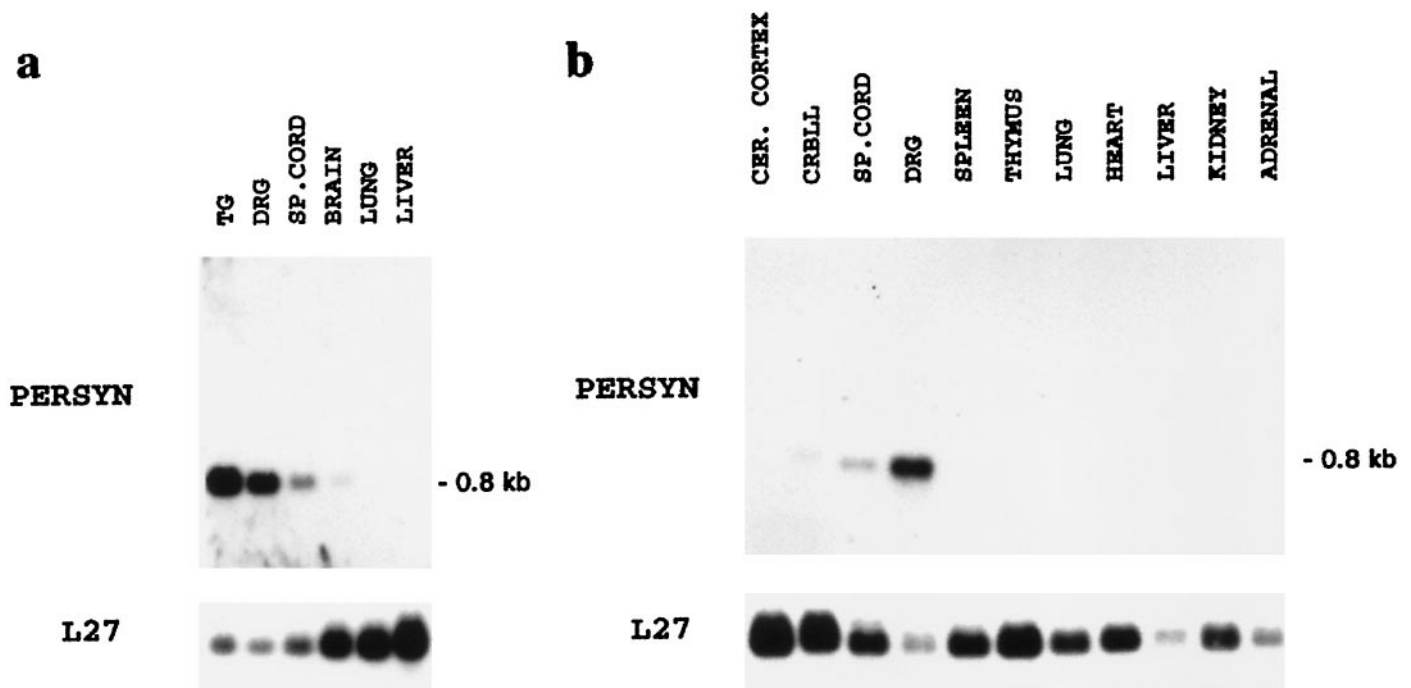


Figure 1. *Persyn* transcripts in mouse and rat tissues. Northern hybridization of 5–20 μ g of total RNA extracted from various tissues of E13 mouse embryos (*a*) and newborn rat (*b*) with a nick-translated *persyn* probe. After stripping off the probe, the same filter was hybridized with a nick-translated cDNA fragment encoding the mouse L27 ribosomal protein to provide an indication of the amount of total RNA from each tissue present on the filter. TG, Trigeminal ganglia; DRG, dorsal root ganglia.

tase, and subtractive hybridization was performed with photobiotinylated poly(A)⁺ RNA (driver). To subtract most of the housekeeping sequences, E13 mouse liver and lung poly(A)⁺ RNAs were used as the driver in two rounds of hybridization. This was followed by two rounds of hybridization with biotinylated poly(A)⁺ RNA from E13 mouse forebrain. cDNAs that failed to form hybrids with biotinylated RNA were cloned as described (Baka and Buchman, 1996). Among the clones isolated from the resulting subtracted cDNA library were two independent overlapped clones, representing *persyn* mRNA.

Miscellaneous cloning procedures. RNA extraction, isolation of poly(A)⁺ RNA, preparation of hybridization probes, Northern, *in situ*, and Southern hybridizations, and library screening were performed as described (Buchman et al., 1992, 1994). Full-length cDNAs were isolated from an E13 mouse trigeminal cDNA library constructed in the λ ZAPII vector and a cDNA library from juvenile human brainstem (Stratagene, La Jolla, CA).

Anti-*persyn* antibody. Rabbits were immunized with the 15-mer C-terminal peptide of mouse *persyn* conjugated with keyhole limpet hemocyanin (Calbiochem, San Diego, CA) activated by MBS (Sigma, St. Louis, MO) (Harlow and Lane, 1988). Monospecific antibodies were purified from the antisera by affinity chromatography using the antigen bound to NHS-activated columns (Supelco, Bellefonte, PA). The anti-mouse monospecific antibody was used at dilutions of 1:500 for Western blot/ECL detection of *persyn* in total cell protein samples. In some experiments 10 ml of diluted antibody was preincubated with 15 μ g of the recombinant mouse *persyn* protein at room temperature for 2 hr.

Protein extraction and Western blotting/ECL detection. Dissected trigeminal ganglia were homogenized directly in SDS-PAGE loading buffer (Laemmli, 1970) and incubated for 5 min in a boiling water bath. Total protein concentrations were measured by the dotMETRIC assay (Geno Technology, St. Louis, MO). Fifteen micrograms of total protein were used for SDS-PAGE (Laemmli, 1970). Electroblooming on Hybond-PVDF membranes was performed in Tris-glycine-methanol buffer as recommended by the membrane supplier (Amersham). Rainbow markers from Amersham were used as protein size standards. After washing with PBS, the membrane was blocked for 1 hr at room temperature in 4% skimmed milk–0.05% Tween 20–PBS. The same buffer was used for incubations with primary and secondary antibodies and washes. The final two washes were in 0.1% Tween 20–PBS. ECL detection was performed as recommended by the Amersham protocol.

Subcellular fractionation. Fractionation of homogenates of the spinal cords, trigeminal ganglia, midbrains, and hindbrains of adult mice was performed as described previously (Cotman and Taylor, 1972; Jones and Matus, 1974). Aliquots of the homogenate (hmg), 1000 gm supernatant (cyt), 14,000 gm supernatant (pmt), 14,000 gm pellet (msk), 120,000 gm supernatant (pmc), 120,000 gm pellet (mcs), mitochondrial (mt), and synaptosomal (syn) fractions of the sucrose gradient were mixed with equal volumes of 2 \times loading buffer (Laemmli, 1970) and incubated for 5 min in a boiling water bath. Twenty micrograms of total protein from each fraction were used for SDS-PAGE followed by Western blotting and detection as described above.

Immunohistochemistry. Fifteen micrometer cryosections were fixed in acetone at -20°C for 10 min and were dried and blocked with 5% goat serum in TBT (20 mM Tris HCl, pH 7.5; 150 mM NaCl; and 0.1% Triton X-100) followed by incubation with the anti-*persyn* antibody (1:50) at 4°C for 16 hr in 1% goat serum–TBT. Sections were washed with 1% goat serum–TBT and processed as recommended by suppliers of secondary HRP-conjugated antibody (Sigma). The same procedure was used for cultured neurons fixed on the Petri dish with cold acetone–methanol mixture (1:1). FITC-conjugated secondary anti-rabbit antibody (Jackson ImmunoResearch, West Grove, PA) was used for detection.

RESULTS

Cloning of *persyn* cDNA

persyn cDNA clones were isolated using the subtractive cloning procedure described in Materials and Methods. Using the insert of the longest clone as a probe for Northern hybridization, a single 0.8 kb transcript was detected only in neural tissues of the E13 mouse embryo. The level of this transcript was much higher in sensory ganglia (trigeminal and dorsal root ganglia) than in CNS, reflecting the way in which the clone was isolated (Fig. 1*a*). Similar results were obtained when the mouse *persyn* probe was used for hybridization with RNAs from newborn rat tissues (Fig. 1*b*).

To isolate a full-length *persyn* cDNA, an E13 mouse trigeminal ganglion cDNA library was constructed and screened with the

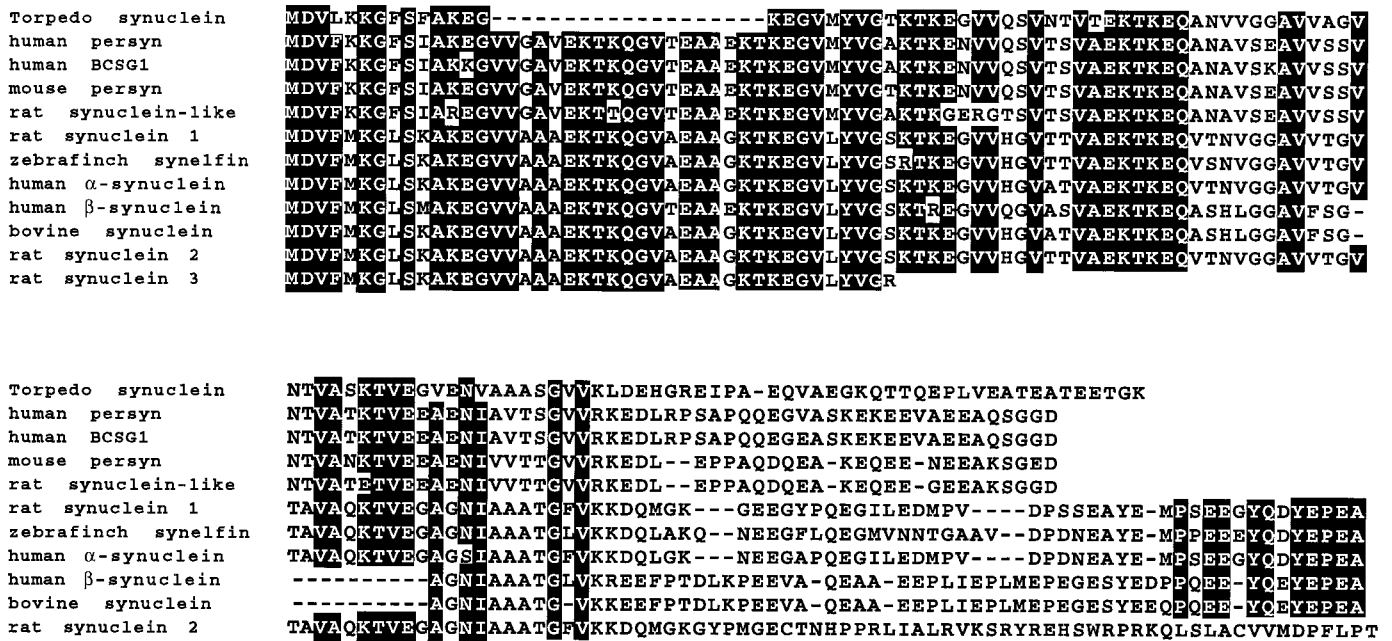


Figure 2. Homologies between mouse persyn and other members of the synuclein family. Alignment of persyn and other members of the synuclein family. Amino acids identical in all or in all but one members of the family are shown in white on black background. Gaps (-) were introduced for better alignment. Sequence accession numbers: Torpedo californica synuclein, P37379; rat synucleins 1, 2, 3, products of alternative splicing of the same gene S73007, S73008, S73009; human α -synuclein (synuclein 1/NACP), L36674; human β -synuclein (synuclein 2), S69965; bovine synuclein, P33567; zebrafish synelfin, L33860; rat synuclein-like, X86789; human BCSG1, AF010126; mouse persyn, AF017255; human persyn, AF017256.

mouse *persyn* cDNA probe. Four overlapping cDNA clones were isolated from this screen, and sequence analysis revealed that the longest of these included 5' untranslated, coding and 3' untranslated (with a characteristic polyadenylation signal) regions (GenBank accession number AF017255). Two human *persyn* cDNA clones were isolated by screening a juvenile brainstem cDNA library with the mouse probe at low stringency. One of these clones included the complete coding region for human persyn protein (GenBank accession number AF017256).

Persyn is a member of the synuclein family

Figure 2 shows the high degree of homology between persyn and other synucleins. All of the EKTKEGV repeats, which are characteristic of the family (Maroteaux et al., 1988; Maroteaux and Scheller, 1991; Uéda et al., 1993; Jakes et al., 1994), are conserved in persyn. The C-terminal region of persyn is highly negatively charged (12 of last 26 amino acids are glutamic or aspartic acid residues), and this region has no obvious homology with other synucleins. The C-terminal amino acid sequences in most synucleins are variable and negatively charged (with the exception of rat synuclein 2, which is charged positively). The sequence of persyn is closely related to the sequence of the founder of the family, Torpedo synuclein (Maroteaux et al., 1988). Two previously published EST sequences that are most closely related to persyn [rat synuclein-like (Akopian and Wood, 1995) and human BCSG1 (Ji et al., 1997)] have multiple substitutions of amino acid that are highly conserved in the synuclein family (Fig. 2). These substitutions are probably resulted from *Taq* polymerase and sequence errors.

***persyn* mRNA is expressed in embryonic sensory and motoneurons**

A digoxigenin-labeled cRNA probe was used for whole-mount *in situ* hybridization of E11 mouse embryos. Consistent with the

results of Northern hybridization, signals were detected in all dorsal root ganglia (DRG) and cranial sensory ganglia and in two strips within the spinal cord (Fig. 3). For precise localization of *persyn*-positive cells, *in situ* hybridization on cryosections was performed. At E10, before the formation of DRG, signals were only detected in the ventral horns of the spinal cord in transverse sections (Fig. 4a). In E11 and E12 embryos, when DRG have become evident, signals were additionally detected in DRG (Fig. 4a). Signals were also observed in all cranial sensory ganglia and in the regions of the hindbrain and midbrain where branchiomotor and somatomotor neurons are located. This is illustrated in the coronal section of an E11 head where hybridization signals in the motoneurons of oculomotor nucleus can be seen (Figs. 4b,c). On transverse sections of postnatal day 9 (P9) mice, *persyn* mRNA was detected in ventral horn motoneurons, most if not all DRG neurons and in the neurons of paravertebral sympathetic chain (Fig. 5a,b).

Persyn in adult brain

Although no persyn protein and mRNA were detected in embryonic and neonatal forebrain structures (Figs. 1, 3, 4) accumulation of persyn protein in the cerebral cortex with age was observed using Western blotting (Buchman et al., 1998). To reveal which kinds of cells express *persyn* mRNA in adult brain we localized *persyn* mRNA by *in situ* hybridization in parasagittal cryosections of rat brain. The strongest hybridization signals were detected in the motoneurons of the brainstem (Fig. 6a). However, substantial levels of expression were found in neurons in many other brain regions including: Purkinje and granule cell layers of the cerebellar cortex and the deep cerebellar nuclei, thalamus, hypothalamus, olfactory bulbs, CA1, CA2, CA3, and CA4 regions of the hippocampus, and all neuron-containing layers of the cerebral cortex in all regions examined (Fig. 6).

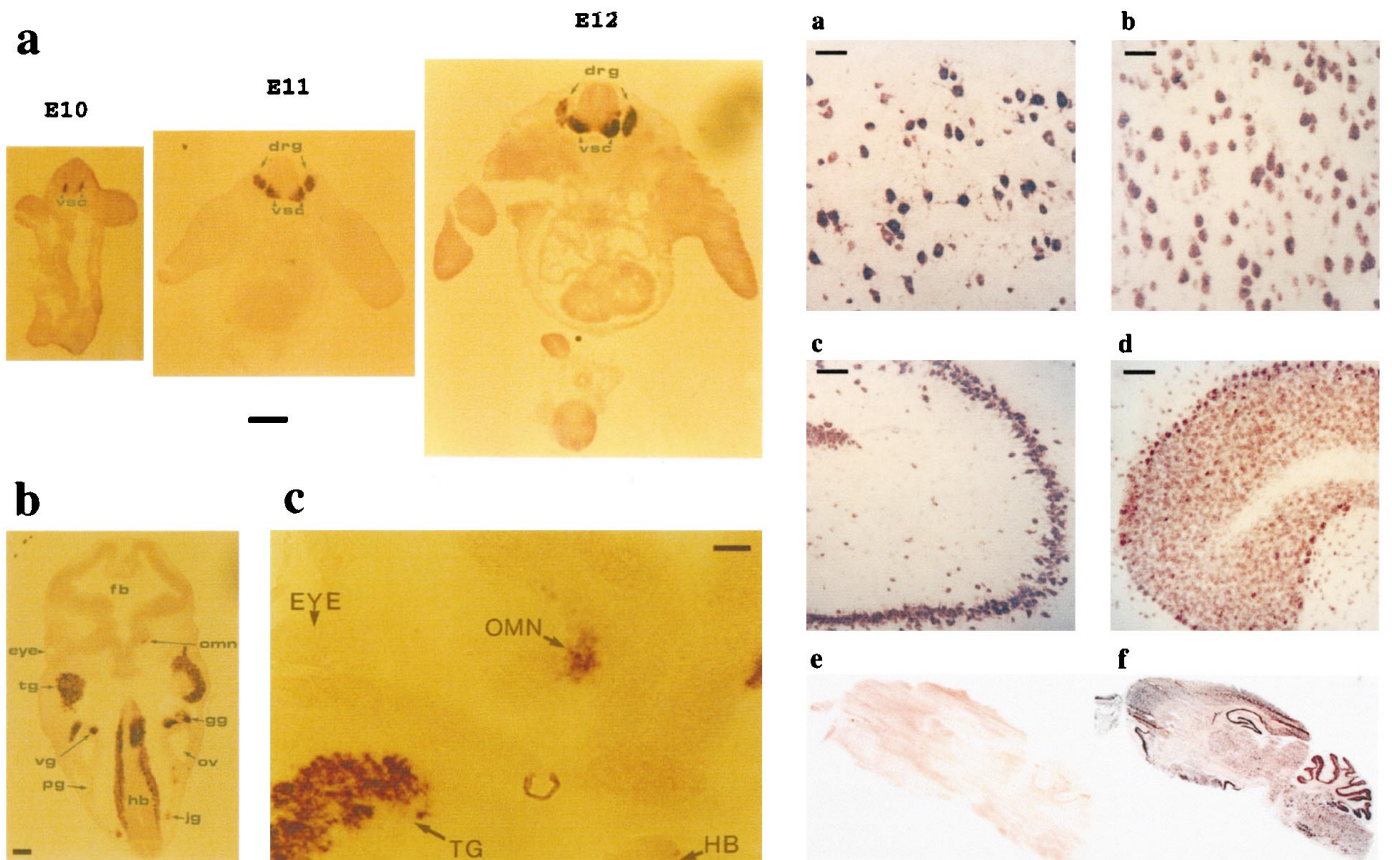
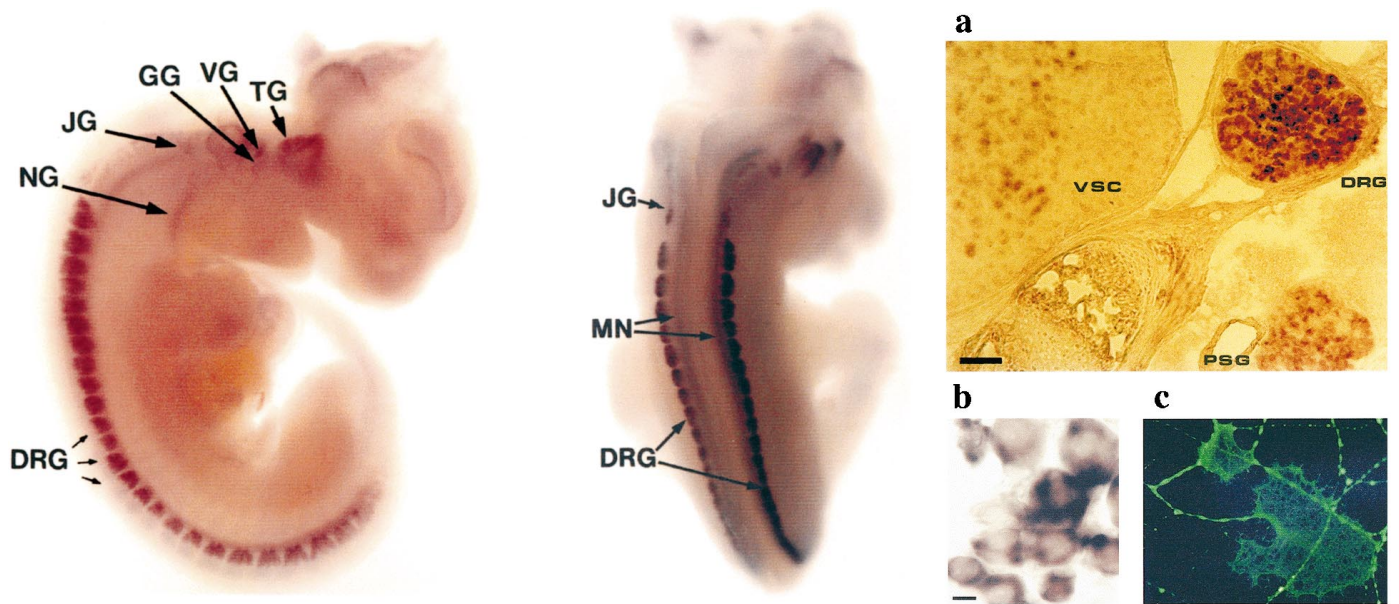


Figure 3. **Top Left.** Whole-mount *in situ* hybridization of mouse embryos. E11 mouse embryos were hybridized with a DIG-labeled antisense cRNA *persyn* probe. *TG*, Trigeminal ganglion; *VG*, vestibular ganglion; *GG*, geniculate ganglion; *JG*, jugular ganglion; *NG*, nodose ganglion; *DRG*, dorsal root ganglia; *MN*, motoneurons in ventral horns of the spinal cord.

Figure 4. **Bottom left.** Detection of *persyn* expression in mouse embryonic tissues by *in situ* hybridization. *a*, Transverse sections at the level of the forelimb buds of *E10*, *E11*, and *E12* embryos were hybridized with a DIG-labeled antisense cRNA *persyn* probe. *vsc*, Motoneurons in ventral horns of the spinal cord; *drg*, dorsal root ganglia. Scale bar, 1 mm. *b*, Coronal section of *E11* mouse embryo head. *tg*, Trigeminal ganglion; *vg*, vestibular ganglion; *gg*, geniculate ganglion; *jg*, jugular ganglion; *pg*, petrosal ganglion; *ov*, otic vesicle; *hb*, hindbrain; *fb*, forebrain; *omn*, oculomotor nucleus. Scale bar, 0.2 mm. *c*, A higher magnification of a coronal section of an *E11* mouse embryo head showing strong labeling in the oculomotor nucleus and trigeminal ganglion. Scale bar, 80 μ m.

Regulation of persyn expression during development of mouse trigeminal ganglion

Because the normal development of the embryonic mouse trigeminal ganglion is known in detail (Davies and Lumsden, 1984, 1986), the expression of persyn was studied in this ganglion at closely staged intervals throughout development to ascertain when persyn is first expressed and how its expression changes during development. Quantitative Northern hybridization revealed that *persyn* mRNA is expressed in the trigeminal ganglion from the earliest stages of its formation. There was a marked increase in expression between E10 and E12, the stage during which the earliest axons are growing to their targets. This high level of expression was maintained into adulthood (Fig. 7*A*). Western blotting revealed that the developmental time course of persyn protein expression in the trigeminal ganglion (Fig. 7*B*) was consistent with the time course of *persyn* mRNA expression observed in Northern blotting studies.

Persyn is cytosolic protein localized in neuronal cell bodies and processes

Specific polyclonal antibodies (SK23) were raised in rabbits against a synthetic C-terminal 15-mer peptide (PS) of mouse persyn. These antibodies detected a single band of ~16 kDa on Western blots that disappeared if the antibodies were preincubated with excess of either the C-terminal peptide or a recombinant persyn protein (Fig. 8*a,b*; data not shown). The mobility of the protein was 3 kDa more than expected from the persyn sequence, but similar decreases in mobility have been reported for other synucleins (Jakes et al., 1994; Weinreb et al., 1996). This antibody does not cross-react with either recombinant α - or β -synucleins and does not detect native α - and β -synucleins in the newborn mouse cerebral cortex (Fig. 8*c*), the tissue where these proteins could be easily detected with specific antibody (Shibayama-Imazu et al., 1993; Hsu et al., 1998).

Western blotting was used to study the distribution of persyn in subcellular fractions of the mouse spinal cord, trigeminal ganglion, midbrain, and hindbrain. Persyn protein was detected only in the cytosolic fractions but not in the particulate and cytoskeletal fractions (Fig. 9). Immunohistochemistry using the SK23 antibodies has shown that persyn is localized in the cell bodies and axons of sensory neurons (Buchman et al., 1998). Likewise, immunocytochemical detection of persyn in cultured DRG neurons showed that persyn is present in the cell bodies, processes, and growth cones of these neurons (Fig. 5*c*) and that it is expressed in both large NT3-dependent and small NGF-dependent neurons (Fig. 5*b*; data not shown).

DISCUSSION

We have cloned and characterized a member of the synuclein family that has a distinctive pattern of expression in the developing and mature nervous system. Persyn shares the main structural features of the family, namely, conservation of all EKTKEGV

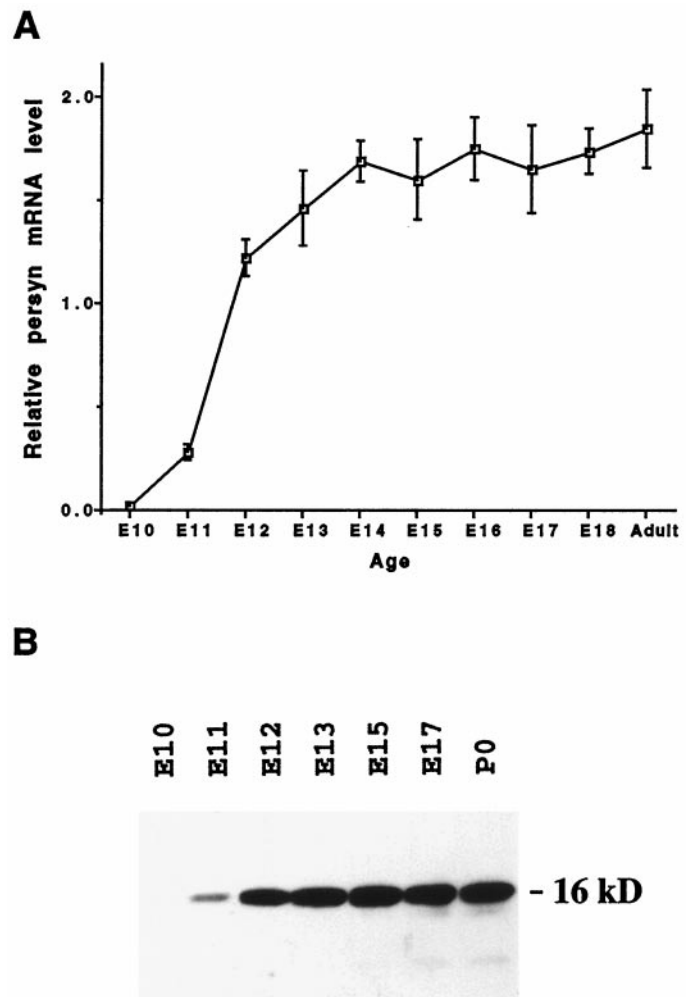


Figure 7. Developmental time course of persyn expression in the trigeminal ganglia. *a*, Graph showing the changes in the level of *persyn* mRNA in the trigeminal ganglia during development. The mean \pm SEM of the persyn mRNA levels relative to GAPDH mRNA levels are shown ($n = 3$). *b*, Western blot showing the developmental changes in the level of persyn protein in the trigeminal ganglia. Equal amounts (15 μ g) of total protein from embryonic (*E*) or newborn (*P0*) mouse trigeminal ganglia were run in each lane.

repeats and the C-terminal part of the molecule is negatively charged. In contrast to α - and β -synucleins, which are predominantly expressed in the cerebral cortex and other forebrain structures (Uéda et al., 1993, 1994; Jakes et al., 1994; George et al., 1995; Iwai et al., 1995a), persyn is abundant in primary sensory neurons and motoneurons. These are the only neurons that express persyn in embryos, although persyn is expressed in sympathetic ganglia in neonates and many different kinds of neurons throughout the brain in the adult.

Figure 5. Top right. Detection of *persyn* expression in mouse postnatal neurons. *a*, *In situ* hybridization of transverse section through the thorax of a P9 mouse neonate with a DIG-labeled antisense cRNA *persyn* probe. *VSC*, Motoneurons of the ventral horns of the spinal cord; *DRG*, dorsal root ganglion; *PSG*, paravertebral sympathetic ganglia. Scale bar, 0.2 mm. *b*, Higher magnification of part of the DRG illustrated in the panel (*a*) showing the cytoplasmic labeling of neurons. *c*, Immunocytochemical localization of persyn protein in processes and growth cones of cultured mouse DRG neurons. Scale bar, 20 μ m.

Figure 6. Bottom right. Detection of *persyn* expression in adult brain by *in situ* hybridization. *In situ* hybridization of parasagittal sections of adult rat brain with a DIG-labeled antisense cRNA *persyn* probe. Labeled neurons within the trigeminal motor nucleus (*a*), frontal cortex (*b*), CA1, and CA3 regions of hippocampus and cerebellar cortex are shown. Scale bars: *a*, *b*, 0.1 mm; *c*, *d*, 0.2 mm. The whole brain images demonstrating absence of hybridization with *persyn* cDNA sense probe (*e*) and specific hybridization with *persyn* cDNA antisense probe (*f*) are shown.

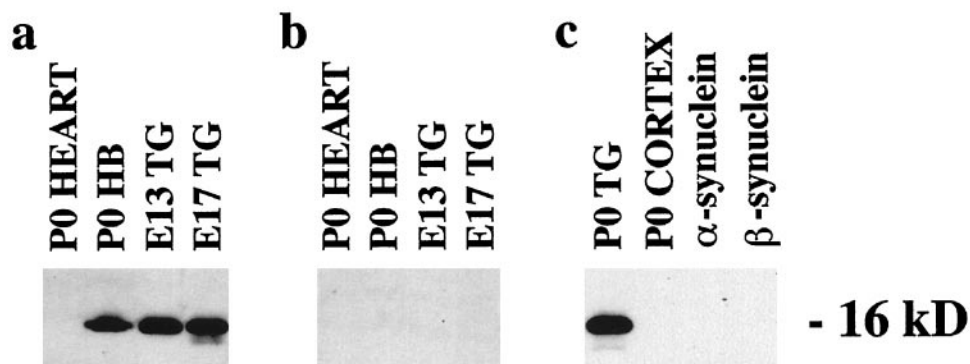
Figure 8. Specificity of anti-mouse persyn antibody. Western blot/ECL detection of persyn protein with affinity-purified SK-23 antibody. *a, b*, Equal amounts (15 μ g) of total protein from P0 mouse heart and hindbrain and E13 and E17 trigeminal ganglia were run in each lane. Western blots were probed with SK-23 antibody (*a*) or SK-23 antibody preincubated with recombinant mouse persyn protein (*b*) as described in Materials and Methods. *c*, Fifteen micrograms of total protein from newborn (P0) mouse trigeminal ganglia (the tissue with a high level of persyn expression) and cerebral cortex (the tissue where persyn expression could not be detected but with substantial levels of α - and β -synucleins expression) and 0.5 μ g of recombinant α - and β -synucleins were analyzed.



Figure 9. Persyn in subcellular fractions from mouse spinal cord. Western blot showing the distribution of persyn protein in subcellular fractions of the adult mouse spinal cord. *hmg*, Homogenate; *cyt*, 1000 gm supernatant; *pmt*, 14,000 gm supernatant; *pmc*, 120,000 gm supernatant; *mcs*, 120,000 gm pellet; *msk*, 14,000 gm pellet; *mt*, mitochondrial fraction; *syn*, synaptosomal fraction. Equal amounts (15 μ g) of total protein from each fraction were run in each lane. Very similar results were obtained with subcellular fractions of trigeminal ganglia, midbrain, and hindbrain.

Immunocytochemical studies suggest that persyn is distributed diffusely throughout the cell body and axons of sensory neurons and motoneurons. This agrees with the results of subcellular fractionation of mouse neuronal tissues that show that persyn is a predominantly cytosolic protein that is not or very loosely associated with cytoskeletal or vesicular fractions. This is similar to the intracellular distribution of human and rat synucleins and zebrafish synelfin (George et al., 1995; Irizarry et al., 1996). However, both α - and β -synuclein are apparently predominantly presynaptic proteins in the cerebral cortex (Iwai et al., 1995a; Masliah et al., 1996; Irizarry et al., 1996).

Although the physiological role of synucleins is unknown, it has been suggested that specific mutations could change the structural and functional properties of these proteins, triggering mechanisms that lead to neurodegeneration (Goedert, 1997; Heintz and Zoghbi, 1997; Nussbaum and Polymeropoulos, 1997; Polymeropoulos et al., 1997). Interestingly, human persyn has a threonine in position 53, which makes it structurally similar to mutated (Ala53Thr) α -synuclein found in some families with hereditary early-onset form of Parkinson's disease (Polymeropoulos et al., 1997). However, what kind of structural and functional consequences this difference could have is not clear. Although it was speculated that Ala53Thr substitution disrupts an α -helix and extends a β -sheet in the predicted structure in this part of the molecule (Polymeropoulos et al., 1997), all experimental attempts to resolve the secondary structure of the wild-type α -synuclein



have led to conclusion that synucleins are “natively unfolded” or random coiled in solution (Weinreb et al., 1996; Kim, 1997). However, it was shown that binding to synthetic membranes stabilizes α -synuclein in an α -helical conformation (Davidson et al., 1998). Perhaps synucleins exist in different conformations in cells depending on their association with membranes, and it is possible that soluble and membrane-bound forms could have different functions.

We have recently shown that persyn plays role in regulating the integrity of the neurofilament network in cultured sensory neurons (Buchman et al., 1998). In this respect, it is interesting that the onset of persyn expression in trigeminal sensory neurons coincides with the stage when axons are starting to grow to their targets and remains high throughout the life, raising the possibility that persyn plays a role in regulating axonal growth and influencing axonal morphology.

The effect of persyn on neurofilaments could be important in neurodegenerative conditions especially given the accumulation of persyn in neurons of the cerebral cortex with age and its localization within axons. Loss of nerve cell processes is believed to contribute significantly to cerebral atrophy in neurodegenerative diseases. Understanding the pathophysiology of this process is important for developing therapeutic strategies to prevent loss of neuronal connectivity. In further work it will be important to investigate the possible contribution of persyn to the loss of neurons and their processes in neurodegenerative diseases.

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